

Sequence alignment of AK110836.q, 83523195.q, 83621446.q, 83621450.q, 83523204.q, 83523203.q, 83523199.q, 83523198.q, 83523201.q, 83523191.q, 83523194.q, 83523187.q, 83500525.q, lir3\_IRK, 2phk\_Phk, and lias\_TGFbR across 1000 amino acids.

The alignment highlights conserved regions in blue, green, and red. Key functional domains include the TGF-β receptor kinase domain (red) and the extracellular matrix binding domain (green).

